Item Listing Version# 1				
Item Name	Download Content			
20100312 095340 us-10-568-055a-91.rag				
20100312 095340 us-10-568-055a-92.rag				
20100312 095341 us-10-568-055a-91.rup				
20100312 095341 us-10-568-055a-92 rup				
<u>20100312 095342 us-10-568-055a-91.rpr</u>				
20100312 095342 us-10-568-055a-92.rpr				
20100312 095343 us-10-568-055a-91 rai				
20100312 095343 us-10-568-055a-92.rai				
20100312 095343 us-10-568-055a-91 rapm				
20100312 095343 us-10-568-055a-92.rapm				
20100312 095343 us-10-568-055a-91.rapn				
20100312 095343 us-10-568-055a-92.rapn				
20100312 095344 us-10-568-055a-91 rapbm				
20100312 095344 us-10-568-055a-92.rapbm				
20100312 095344 us-10-568-055a-91 rapbn				
20100312 095344 us-10-568-055a-92.rapbn				

RESULT 1

US-10-568-055A-92

- ; Sequence 92, Application US/10568055A
- ; GENERAL INFORMATION
- ; APPLICANT: Cambridge University Technical Services Limited
- ; APPLICANT: Doherty, Aidan
- ; APPLICANT: Della, Marina
- ; APPLICANT: Weller, Geoffrey
- ; APPLICANT: Jackson, Stephen
- ; TITLE OF INVENTION: Prokaryotic DNA Repair Ligases
- ; FILE REFERENCE: 6947-73362-01
- ; CURRENT APPLICATION NUMBER: US/10/568,055A
- ; CURRENT FILING DATE: 2006-09-27
- ; PRIOR APPLICATION NUMBER: PCT/GB2004/003349
- ; PRIOR FILING DATE: 2004-08-02
- ; PRIOR APPLICATION NUMBER: US 60/494,088
- ; PRIOR FILING DATE: 2003-08-12
- ; NUMBER OF SEQ ID NOS: 92
- ; SOFTWARE: PatentIn version 3.1
- ; SEQ ID NO 92
- ; LENGTH: 273
- ; TYPE: PRT

```
; ORGANISM: Mycobacterium tuberculosis
US-10-568-055A-92
 Query Match
                     100.0%; Score 1405; DB 5; Length 273;
 Best Local Similarity 100.0%;
 Matches 273; Conservative 0; Mismatches
                                         0;
                                            Indels
                                                    0; Gaps
0:
         1 MRAIWTGSIAFGLVNVPVKVYSATADHDIRFHQVHAKDNGRIRYKRVCEACGEVVDYRDL 60
Qу
           Db
         1 MRAIWTGSIAFGLVNVPVKVYSATADHDIRFHOVHAKDNGRIRYKRVCEACGEVVDYRDL 60
         61 ARAYESGDGQMVAITDDDIASLPEERSREIEVLEFVPAADVDPMMFDRSYFLEPDSKSSK
Qy
120
           Db
         61 ARAYESGDGQMVAITDDDIASLPEERSREIEVLEFVPAADVDPMMFDRSYFLEPDSKSSK
120
        121 SYVLLAKTLAETDRMAIVHFTLRNKTRLAALRVKDFGKREVMMVHTLLWPDEIRDPDFPV
Qу
180
           121 SYVLLAKTLAETDRMAIVHFTLRNKTRLAALRVKDFGKREVMMVHTLLWPDEIRDPDFPV
Dh
180
        181 LDOKVEIKPAELKMAGOVVDSMADDFNPDRYHDTYOEOLOELIDTKLEGGOAFTAEDOPR
Qу
240
           181 LDQKVEIKPAELKMAGQVVDSMADDFNPDRYHDTYQEQLQELIDTKLEGGQAFTAEDQPR
Db
240
        241 LLDEPEDVSDLLAKLEASVKARSKANSNVPTPP 273
Qу
           241 LLDEPEDVSDLLAKLEASVKARSKANSNVPTPP 273
Db
EAST SEARCH:
BRS
    L4
         1062 ligase and helicase
                                 USPAT 2011/05/20 15:40
    L5
         18
              (ligase and helicase).clm.
                                      USPAT 2011/05/20 15:40
BRS
STN SEARCH:
 FILE 'MEDLINE, BIOSIS, BIOTECHNO, CAPLUS, EMBASE, JAPIO' ENTERED AT
   16:09:41 ON 20 MAY 2011
         887 S HELICASE AND LIGASE
L2
ь3
          85 S L2 AND METHOD
```

80 DUP REM L3 (5 DUPLICATES REMOVED)

L4